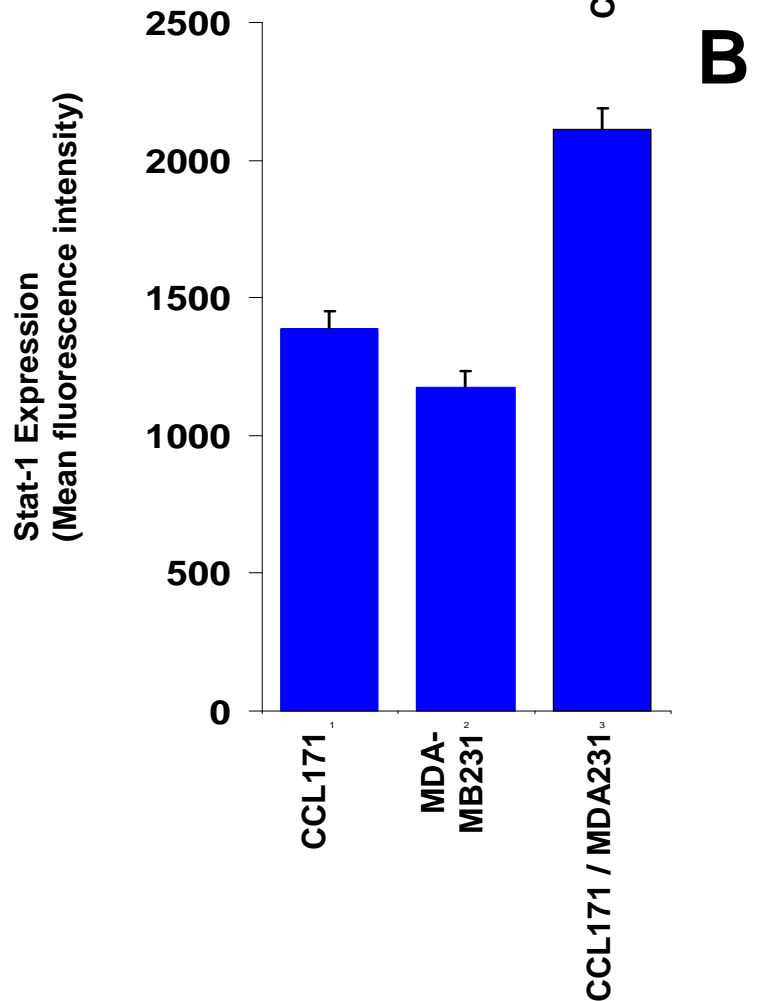
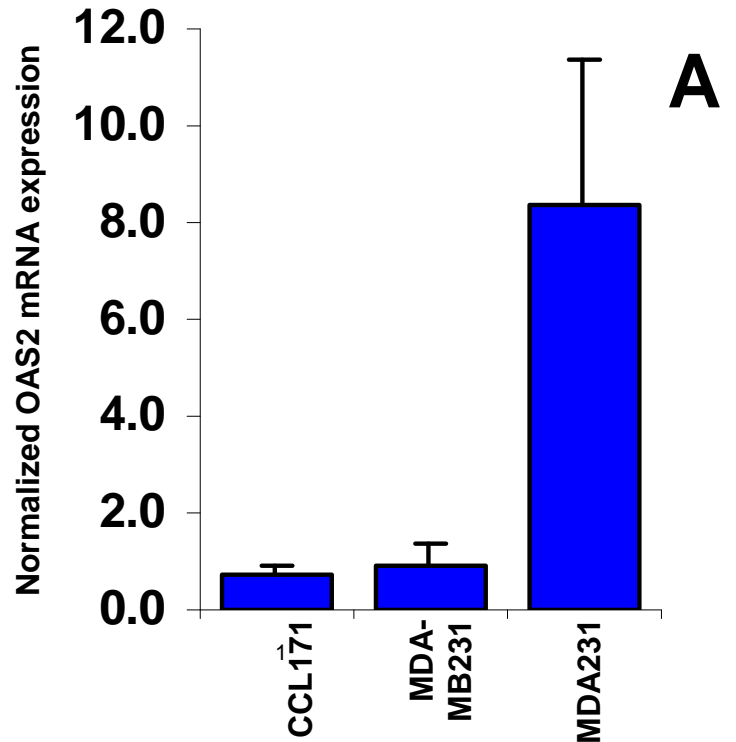


## Additional File 2

### Legend Figure A:

Expression of OAS2 was determined by quantitative RT-PCR. Expression of GAPDH was used for normalization between the samples. OAS2 was up-regulated in CCL171/MDA-MB-231 co-culture compared to these cells cultivated as monocultures.



### Legend Figure B:

Expression of STAT1 determined by immunofluorescent staining and FACS analysis.